
Sequence Listing was accepted.

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Reviewer: Durreshwar Anjum

Timestamp: Wed Oct 24 13:27:30 EDT 2007

Validated By CRFValidator v 1.0.3

Application No: 10511794 Version No: 2.0

0

Input Set:

Output Set:

Started: 2007-10-05 15:51:15.002 **Finished:** 2007-10-05 15:51:16.476

Elapsed: 0 hr(s) 0 min(s) 1 sec(s) 474 ms

Total Warnings: 23

No. of SeqIDs Defined: 23

Actual SeqID Count: 23

Total Errors:

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Input Set:

Output Set:

Started: 2007-10-05 15:51:15.002 **Finished:** 2007-10-05 15:51:16.476

Elapsed: 0 hr(s) 0 min(s) 1 sec(s) 474 ms

Total Warnings: 23
Total Errors: 0

No. of SeqIDs Defined: 23

Actual SeqID Count: 23

Error code Error Description

This error has occured more than 20 times, will not be displayed

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<151> 2003-04-28
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Ala Phe Ile Ser Ser Asp Gly Ile Ala Tyr Tyr Ala Asp Ser Val Lys
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50

55

60

Gly 65	Arg	Phe	Thr	Ile	Ser 70	Arg	Asp	Asn	Ala	Arg 75	Asn	Ile	Leu	Tyr	Leu 80
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Arg	Val	Tyr	Tyr 100	Tyr	Gly	Ser	Ser	Tyr 105	Phe	Asp	Tyr	Trp	Gly 110	Gln	Gly
Thr	Thr	Leu 115	Thr	Val	Ser	Ser	Glu 120	Gly	Lys	Ser	Ser	Gly 125	Ser	Gly	Ser
Glu	Ser 130	Lys	Val	Asp	Asp	Ile 135	Val	Met	Thr	Gln	Ser 140	Pro	Lys	Phe	Met
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Asn	Ala	Gly	Thr	Asn 165	Val	Ala	Trp	Tyr	Gln 170	Gln	Lys	Pro	Gly	Gln 175	Ser
Pro	Lys	Ala	Leu 180	Ile	Tyr	Ser	Ala	Ser 185	Ser	Arg	Asn	Ser	Gly 190	Val	Pro
Asp	Arg	Ile 195	Thr	Gly	Ser	Gly	Ser 200	Gly	Thr	Asp	Phe	Thr 205	Leu	Thr	Ile
Ser	Asn 210	Val	Gln	Ser	Glu	Asp 215	Leu	Ala	Glu	Tyr	Phe 220	Суз	Gln	Gln	Tyr
Asn 225	Ser	Tyr	Pro	Leu	Val 230	Thr	Phe	Gly	Ala	Gly 235	Thr	Lys	Leu	Glu	Leu 240
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Ala	Met	Ser 35	Trp	Val	Arg	Gln	Thr 40	Pro	Glu	Lys	Arg	Leu 45	Glu	Trp	Val
7. 7	D.1	- 1	a	a	-	a i	- 1	3.7			3.7		a		-

Ala Phe Ile Ser Ser Asp Gly Ile Ala Tyr Tyr Ala Asp Ser Val Lys

Gly Arg Phe Thr Ile Ser Arg Asp Asn Ala Arg Asn Ile Leu Tyr Leu Gln Met Ser Ser Leu Arg Ser Glu Asp Thr Ala Met Tyr Tyr Cys Ala 8.5 90 Arg Val Tyr Tyr Gly Ser Ser Tyr Phe Asp Tyr Trp Gly Gln Gly 105 Thr Thr Leu Thr Val Ser Ser Gly Gly Gly Ser Asp Ile Val Met 115 120 Thr Gln Ser Pro Lys Phe Met Ser Thr Ser Val Gly Asp Arg Val Ser 135 Val Thr Cys Lys Ala Ser Gln Asn Ala Gly Thr Asn Val Ala Trp Tyr 145 150 155 Gln Gln Lys Pro Gly Gln Ser Pro Lys Ala Leu Ile Tyr Ser Ala Ser 165 170 Ser Arg Asn Ser Gly Val Pro Asp Arg Ile Thr Gly Ser Gly Ser Gly 180 185 190 Thr Asp Phe Thr Leu Thr Ile Ser Asn Val Gln Ser Glu Asp Leu Ala 200 195 Glu Tyr Phe Cys Gln Gln Tyr Asn Ser Tyr Pro Leu Val Thr Phe Gly 215 Ala Gly Thr Lys Leu Glu Leu Lys 225 230 <210> 18 <211> 35 <212> DNA <213> Artificial Sequence <220> <223> Description of Artificial Sequence: oligo <400> 18 catgccatgg ggaatccgaa gtgaagctgg tggag 35 <210> 19 <211> 36 <212> DNA <213> Artificial Sequence <220> <223> Description of Artificial Sequence: oligo <400> 19

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                            40
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Gly Arg Phe Thr Ile Ser Arg Asp Asn Ala Arg Asn Ile Leu Tyr Leu
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Arg Val Tyr Tyr Gly Ser Ser Tyr Phe Asp Tyr Trp Gly Gln Gly
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           100
Thr Thr Leu Thr Val Ser Ser Gly Gly Gly Ser Asp Ile Ile Met
       115
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                                          140
Val Thr Cys Lys Ala Ser Gln Asn Ala Gly Thr Asn Val Ala Trp Tyr
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Gln Gln Lys Pro Gly Gln Ser Pro Lys Ala Leu Ile Tyr Ser Ala Ser
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Ser Arg Asn Ser Gly Val Pro Asp Arg Ile Thr Gly Ser Gly Ser Gly

180 185 190

Thr Asp Phe Thr Leu Thr Ile Ser Asn Val Gln Ser Glu Asp Leu Ala 200 Glu Tyr Phe Cys Gln Gln Tyr Asn Ser Tyr Pro Leu Val Thr Phe Gly 215 220 Ala Gly Thr Lys Leu Glu Leu Lys Ala Ala Gly Ser Glu Gln Lys 230 235 Leu Ile Ser Glu Glu Asp Leu Asn Ser His His His His His 245 250 <210> 22 <211> 4 <212> PRT <213> Artificial Sequence <220> <223> Synthetic amino acid sequence <220> <221> DISULFID <222> (2)..(3) <400> 22 Phe Arg Ser Arg <210> 23 <211> 4 <212> PRT <213> Artificial Sequence <220> <223> Synthetic amino acid sequence <220> <221> DISULFID <222> (2)..(3) <400> 23

Val Lys Ile Lys